Validated By CRFValidator v 1.0.3

Application No: 10539402 Version No: 2.0

Input Set:

Output Set:

Started: 2009-09-01 16:50:17.498
Finished: 2009-09-01 16:50:21.960
Elapsed: 0 hr(s) 0 min(s) 4 sec(s) 462 ms
Total Warnings: 60
Total Errors: 0
No. of SeqIDs Defined: 162
Actual SeqID Count: 162

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (73)
W 402	Undefined organism found in <213> in SEQ ID (74)
W 213	Artificial or Unknown found in <213> in SEQ ID (109)
W 213	Artificial or Unknown found in <213> in SEQ ID (110)
W 213	Artificial or Unknown found in <213> in SEQ ID (111)
W 213	Artificial or Unknown found in <213> in SEQ ID (112)
W 213	Artificial or Unknown found in <213> in SEQ ID (113)
W 213	Artificial or Unknown found in <213> in SEQ ID (114)
W 213	Artificial or Unknown found in <213> in SEQ ID (115)
W 213	Artificial or Unknown found in <213> in SEQ ID (116)
W 213	Artificial or Unknown found in <213> in SEQ ID (117)
W 213	Artificial or Unknown found in <213> in SEQ ID (118)
W 213	Artificial or Unknown found in <213> in SEQ ID (119)
W 213	Artificial or Unknown found in <213> in SEQ ID (120)
W 213	Artificial or Unknown found in <213> in SEQ ID (121)
W 213	Artificial or Unknown found in <213> in SEQ ID (122)

Input Set:

Output Set:

Started: 2009-09-01 16:50:17.498
Finished: 2009-09-01 16:50:21.960
Elapsed: 0 hr(s) 0 min(s) 4 sec(s) 462 ms
Total Warnings: 60
Total Errors: 0
No. of SeqIDs Defined: 162
Actual SeqID Count: 162

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (123)
W 213	Artificial or Unknown found in <213> in SEQ ID (124)
W 213	Artificial or Unknown found in <213> in SEQ ID (125)
W 213	Artificial or Unknown found in <213> in SEQ ID (126)
W 213	Artificial or Unknown found in <213> in SEQ ID (127)
W 213	Artificial or Unknown found in <213> in SEQ ID (128)

This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

<110> UNGER, CHRISTINE MARGARETE
BESTE, GERALD
ZEHETMEIER, CAROLIN
LAIN, BLANCA
TORELLA, CLAUDIA
NIEWOHNER, JENS
JAY, DANIEL G.
EUSTACE, BRENDA K.
KNAUER, ROLAND
JENSEN, KRISTIAN HOBOLD

<120> NEUROPILIN-1 INHIBITORS

<130> MXI-352US

<140> 10539402
<141> 2005-12-22

<150> PCT/EP2003/014756
<151> 2003-12-22

<150> EP 03000615.9
<151> 2003-01-15

<150> 60/435,893
<151> 2002-12-20

<160> 162

<170> PatentIn version 3.5

<210> 1
<211> 269
<212> PRT
<213> Mus sp.

<400> 1

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Glu Val Gln Leu Gln Gln Ser Gly Pro Glu Leu Val Lys Pro Gly Ala
1               5                   10                  15


Leu Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr Thr Val Thr Ser Tyr
            20                  25                  30


Asp Ile Asn Trp Val Lys Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile
        35                  40                  45


Gly Trp Ile Tyr Pro Gly Asp Gly Ser Thr Lys Tyr Asn Glu Lys Phe
    50                  55                  60


Lys Gly Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Thr Thr Val Tyr
```

```
65                  70                  75                  80

Met Gln Leu Ser Ser Leu Thr Ser Glu Asn Ser Ala Val Tyr Phe Cys
                85                  90                  95

Ala Arg Gly Gly Lys Tyr Phe Asp Tyr Trp Gly Gln Gly Thr Thr Leu
            100                 105                 110

Thr Val Ser Thr Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly
        115                 120                 125

Gly Gly Ser Ala Leu Asp Ile Val Met Thr Gln Ser Pro Lys Phe Met
    130                 135                 140

Ser Thr Ser Val Gly Asp Arg Val Ser Val Thr Cys Lys Ala Ser Gln
145                 150                 155                 160

Asn Val Ala Thr Asn Val Ala Trp Tyr Gln Gln Lys Pro Gly Gln Ser
                165                 170                 175

Pro Lys Pro Leu Thr Tyr Ser Ala Ser Phe Arg Ser Ser Gly Val Pro
            180                 185                 190

Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile
        195                 200                 205

Ser Asn Val Gln Ser Glu Asp Leu Ala Glu Tyr Phe Cys Gln Gln Tyr
    210                 215                 220

Asn Ser Tyr Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys
225                 230                 235                 240

Ala Ala Ala Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu Glu Pro Arg
                245                 250                 255

Gly Ala Ala Ser Ala Trp Ser His Pro Gln Phe Glu Lys
            260                 265
```

<210> 2
<211> 288
<212> PRT
<213> Mus sp.

<400> 2

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Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
1               5                  10                  15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
            20                  25                  30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
                85                  90                  95

Ala Arg Asp Ser Gly Leu Gln Gln Gly Pro Arg Arg Arg Gly Ala Arg
            100                 105                 110

Val Asn Phe Ser Tyr Tyr Gly Leu Asp Val Trp Gly Arg Gly Thr Thr
        115                 120                 125

Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
    130                 135                 140

Gly Gly Gly Ser Ala Gln Ala Val Leu Thr Gln Pro Ser Ser Ala Ser
145                 150                 155                 160

Gly Thr Pro Gly Gln Arg Val Thr Ile Ser Cys Ser Gly Ser Asn Ser
                165                 170                 175

Asn Ile Gly Arg Asn Tyr Val Phe Trp Tyr Gln Gln Phe Pro Gly Thr
            180                 185                 190

Ala Pro Lys Ile Leu Ile Tyr Arg Asn Asn Gln Arg Pro Ser Gly Val
        195                 200                 205

Pro Asp Arg Phe Ser Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala
    210                 215                 220

Ile Ser Gly Leu Arg Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ser
```

225 230 235 240

Trp Asp Asp Ser Leu Thr Trp Val Phe Gly Gly Gly Thr Lys Val Thr
245 250 255

Val Leu Gly Ala Ala Ala Gly Ala Pro Val Pro Tyr Pro Asp Pro Leu
260 265 270

Glu Pro Arg Gly Ala Ala Ser Ala Trp Ser His Pro Gln Phe Glu Lys
275 280 285

<210> 3
<211> 810
<212> DNA
<213> Mus sp.

<400> 3
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tcctgcaagg cctcgggata caccgtcaca agctacgata taaactgggt gaagcagagg 120

cctggacagg gacttgagtg gattggatgg atttatcctg gagatggtag tactaagtac 180

aatgagaaat tcaagggcaa ggccacactg actgtagaca aatcctccac cacagtctac 240

atgcagctca gcagcctgac ttctgagaac tctgcagtct atttctgtgc aagaggtggt 300

aaatactttg actactgggg ccaaggcacc actctcacag tgtcgacagg tggaggcggt 360

tcaggcggag gtggctctgg cggtggcgga agtgcactcg acattgtgat gacacagtct 420

ccaaaattca tgtccacatc agtaggagac agggtcagcg tcacctgcaa ggccagtcag 480

aatgtggcta ctaatgtagc ctggtatcaa cagaaaccag ggcaatctcc taaaccactg 540

acttactcgg catccttccg gtccagtgga gtccctgatc gcttcacagg cagtggatct 600

gggacagatt tcactctcac catcagcaat gtgcagtctg aagacttggc agagtatttc 660

tgtcagcaat ataacagcta tccgtacacg ttcggagggg ggaccaagct ggaaataaaa 720

gcggccgcag gtgcgccggt gccgtatcca gatccgctgg aaccgcgtgg ggccgcaagc 780

gcttggagcc acccgcagtt cgaaaaataa 810

<210> 4
<211> 867
<212> DNA
<213> Mus sp.

<400> 4
gaggtgcagc tgttggagtc tgggggaggc ttggtacagc ctgggggggtc cctgagactc 60

tcctgtgcag cctctggatt cacctttagc agctatgcca tgagctgggt ccgccaggct 120

ccagggaagg ggctggagtg ggtctcagct attagtggta gtggtggtag cacatactac 180

gcagactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacgctgtat 240

ctgcaaatga acagcctgag agccgaggac acggccgtgt attactgtgc gcgagactcg 300

gggctacagc agggaccccg ccgaagaggg gcccgagtaa atttctccta ctacggtctg 360

gacgtctggg ggcgggggac cacggtcacc gtctcgagtg gaggcggcgg ttcaggcgga 420

ggtggctctg gcggtggcgg aagtgcacag gctgtgctga ctcagccgtc ctcagcgtct 480

gggacccccg ggcagagggt caccatctct tgttctggaa gcaactccaa catcggacgc 540

aattatgtat tctggtacca gcagttccca ggaacggccc ccaaaatcct catctacagg 600

aacaatcagc ggccctcagg ggtccctgac cgattctctg gctccaagtc tggcacatca 660

gcctccctgg ccatcagtgg gctccggtcc gaggatgagg ctgattatta ctgtgcatca 720

tgggatgaca gcctgacttg ggtgttcggc ggagggacca aggtcaccgt cctaggtgcg 780

gccgcaggtg cgccggtgcc gtatccagat ccgctggaac cgcgtggggc cgcaagcgct 840

tggagccacc cgcagttcga aaaataa 867

<210> 5
<211> 246
<212> PRT
<213> Homo sapiens

<400> 5

Ala	Ser	Val	Lys	Val	Ser	Cys	Lys	Thr	Ser	Gly	Tyr	Thr	Phe	Ile	Ala
1				5				10					15		

Tyr	Tyr	Ile	His	Trp	Val	Arg	Gln	Ala	Pro	Gly	Gln	Gly	Leu	Glu	Trp
			20					25					30		

Met	Gly	Arg	Ile	Asn	Pro	Asn	Thr	Gly	Gly	Ile	Asn	Leu	Ala	Gln	Lys
		35					40					45			

Phe	Gln	Gly	Arg	Val	Thr	Val	Thr	Arg	Asp	Thr	Ser	Ile	Ser	Thr	Ala
	50					55					60				

His	Met	Glu	Leu	Ser	Arg	Leu	Ser	Ser	Asp	Asp	Thr	Ala	Val	Tyr	Tyr
65					70					75					80

Cys	Ala	Arg	Glu	Arg	Ile	Val	Pro	Ala	Gly	Leu	Arg	Asn	Arg	Gly	Met

```
                85                  90                  95

Val Thr Ala Val Gly Met Asp Val Trp Gly Arg Gly Thr Leu Val Thr
            100                 105                 110

Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly
        115                 120                 125

Gly Ser Ala Gln Ser Val Val Thr Gln Pro Pro Ser Met Ser Gly Thr
    130                 135                 140

Pro Gly Gln Arg Val Thr Ile Ser Cys Ser Gly Ser Arg Ser Asn Ile
145                 150                 155                 160

Gly Arg Asn Tyr Val Tyr Trp Tyr Gln Gln Phe Pro Gly Thr Ala Pro
                165                 170                 175

Lys Leu Leu Ile Tyr Arg Asn Asn Glu Arg Pro Ser Gly Val Pro Asp
            180                 185                 190

Arg Phe Ser Ala Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser
        195                 200                 205

Gly Leu Arg Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Thr Trp Asp
    210                 215                 220

Asp Ser Leu Ser Gly Thr Trp Val Phe Gly Gly Gly Thr Lys Leu Thr
225                 230                 235                 240

Val Leu Gly Ala Ala Ala
                245
```

<210> 6
<211> 248
<212> PRT
<213> Homo sapiens

<400> 6

```
Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly Ser Leu Arg
1               5                   10                  15

Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr Ala Met Ser
            20                  25                  30
```

```
Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ser Ala Ile
        35                  40                  45

Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg
    50                  55                  60

Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln Met
65                  70                  75                  80

Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg Gly
                85                  90                  95

Gly Gly Arg Tyr Asp Ser Ser His Gly Phe Asp Ser Trp Gly Arg Gly
            100                 105                 110

Thr Met Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly
        115                 120                 125

Ser Gly Gly Gly Gly Ser Ala Leu Ser Tyr Glu Leu Thr Gln Pro Pro
    130                 135                 140

Ser Val Ser Val Ala Pro Gly Glu Thr Ala Thr Ile Thr Cys Gly Gly
145                 150                 155                 160

Arg Ser Leu Gly Ser Lys Val Val His Trp Tyr Gln Gln Lys Pro Gly
                165                 170                 175

Gln Ala Pro Thr Leu Val Ile Tyr Tyr Asp Ser Val Arg Pro Ser Gly
            180                 185                 190

Val Pro Glu Arg Phe Ser Ala Ser Asn Ser Arg Leu Ser Ala Thr Leu
        195                 200                 205

Thr Val Ser Arg Val Glu Ala Gly Asp Glu Ala Asp Tyr Tyr Cys Gln
    210                 215                 220

Val Trp Asp Arg Ser Ser Asp His Tyr Val Phe Gly Thr Gly Thr Lys
225                 230                 235                 240

Leu Thr Val Leu Gly Ala Ala Ala
                245
```

<210> 7

<211> 248
<212> PRT
<213> Homo sapiens

<400> 7
Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly Ser Leu
1   5   10   15

Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr Ala Met
            20   25   30

Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ser Ala
        35   40   45

Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys Gly
    50   55   60

Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln
65   70   75   80

Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg
                85   90   95

Asp Trp Arg Trp Gln Gln Phe Gly Gly Trp Phe Asp Pro Trp Gly Arg
            100   105   110

Gly Thr Leu Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly
        115   120   125

Gly Ser Gly Gly Gly Gly Ser Ala Leu Glu Thr Thr Leu Thr Gln Ser
    130   135   140

Pro Ala Thr Leu Ser Leu Ser Pro Gly Glu Thr Ala Thr Leu Phe Cys
145   150   155   160

Arg Ala Ser Gln Ser Val Arg Asn Asn Leu Val Trp Tyr Gln Gln Lys
                165   170   175

Leu Gly Gln Ala Pro Arg Leu Leu Ile Phe Gly Ala Ser Thr Arg Ala
            180   185   190

Ser Gly Ile Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe
        195   200   205

```
Ser Leu Thr Ile Thr Lys Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr
    210                 215                 220

Cys Gln Arg Tyr Gly Gly Phe Pro Ile Thr Phe Gly Gln Gly Thr Arg
225                 230                 235                 240

Leu Glu Ile Lys Arg Ala Ala Ala
                245
```

<210> 8
<211> 247
<212> PRT
<213> Homo sapiens

<400> 8

```
Gln Leu Val Gln Ser Gly Gly Gly Leu Val Gln Pro Gly Gly Ser Leu
1               5                   10                  15

Arg Leu Ala Cys Glu Ala Ser Gly Phe Arg Phe Ser Ser Tyr Gly Met
            20                  25                  30

Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val Ser Ser
        35                  40                  45

Met Ser Asp Ser Gly Ala Asn Thr Tyr Tyr Ala Asp Ser Val Lys Gly
    50                  55                  60

Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Lys Met Leu Tyr Leu Gln
65                  70                  75                  80

Met Ser Ser Leu Arg Gly Glu Asp Thr Ala Val Tyr Tyr Cys Ala Thr
                85                  90                  95

Leu Phe Arg Gly Ser Gly Tyr Val Arg His Trp Gly Arg Gly Thr Leu
            100                 105                 110

Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
        115                 120                 125

Gly Gly Gly Ser Ala Gln Ala Val Leu Thr Gln Pro Ser Ser Ala Ser
    130                 135                 140

Gly Thr Pro Gly Gln Arg Val Ile Ile Ser Cys Ser Gly Ser Ser Ser
145                 150                 155                 160
```

```
Asn Ile Ala Ser Asn Tyr Val Tyr Trp Tyr Gln Gln Leu Pro Gly Thr
                165                 170                 175

Ala Pro Lys Leu Leu Ile Ser Lys Asn Ser Arg Arg Pro Ser Gly Val
            180                 185                 190

Pro Asp Arg Phe Ser Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala
        195                 200                 205

Ile Ser Glu Leu Arg Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala
    210                 215                 220

Trp Asp Asp Arg Leu Ser Gly Pro Ala Phe Gly Gly Gly Thr Lys Leu
225                 230                 235                 240

Thr Val Leu Gly Ala Ala Ala
                245
```

<210> 9
<211> 248
<212> PRT
<213> Homo sapiens

<400> 9

```
Lys Lys Pro Gly Ser Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly
1               5                   10                  15

Thr Phe Ser Ser Tyr Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln
            20                  25                  30

Gly Leu Glu Trp Met Gly Gly Ile Ile Pro Met Ser Gly Thr Pro Asn
        35                  40                  45

Tyr Ala Gln Lys Phe Gln Asp Arg Val Thr Ile Thr Ala Asp Lys Ser
    50                  55                  60

Thr Ser Thr Ala Tyr Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr
65                  70                  75                  80

Ala Val Tyr Tyr Cys Ala Arg Gly Gly Arg Tyr Val Asp Phe Gly Arg
                85                  90                  95

Gly Pro Ser Tyr His Tyr Tyr Tyr Met Asp Val Trp Gly Arg Gly Thr
            100                 105                 110
```

```
Leu Val Thr Val Ser Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
        115                 120                 125

Gly Gly Gly Gly Ser Ala Gln Ser Val Leu Thr Gln Pro Pro Ser Ala
    130                 135                 140

Ser Gly Thr Pro Gly Gln Arg Val Thr Ile Ser Cys Ser Gly Ala Thr
145                 150                 155                 160

Ser Asn Ile Gly Arg Asn Tyr Val Tyr Trp Tyr His Gln Leu Pro Gly
                16
```